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## SEQUENCE LISTING

COPY OF PAPERS  
ORIGINALLY FILED<110> LAZDUNSKI, MICHEL  
LAMBEAU, GERARD  
VALENTIN, EMMANUEL<120> CLONING AND RECOMBINANT EXPRESSION OF MAMMALIAN GROUP  
XII SECRETED PHOSPHOLIPASE A2

&lt;130&gt; 1479-R-00

&lt;140&gt; 09/975,374

&lt;141&gt; 2001-10-11

&lt;150&gt; 60/239,489

&lt;151&gt; 2000-10-11

&lt;160&gt; 18

&lt;170&gt; PatentIn Ver. 2.1

&lt;210&gt; 1

&lt;211&gt; 716

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (121)..(690)

&lt;223&gt; cDNA coding the human group XII sPLA2

&lt;400&gt; 1

atatggagct ggctgctgcc aagtccgggg cccgcgccgc tgcctagcgc gtcctgggga 60

ctctgtgggg acgcgccccg cgccgcgggt cggggaccgc tagagcccg cgtgcgcgc 120

atg gcc ctg ctc tcg cgc ccc gcg ctc acc ctc ctg ctc ctc ctc atg 168  
Met Ala Leu Leu Ser Arg Pro Ala Leu Thr Leu Leu Leu Leu Met  
1 5 10 15gcc gct gtt gtc agg tgc cag gag cag gcc cag acc acc gac tgg aga 216  
Ala Ala Val Val Arg Cys Gln Glu Gln Ala Gln Thr Thr Asp Trp Arg  
20 25 30gcc acc ctg aag acc atc cgg aac ggc gtt cat aag ata gac acg tac 264  
Ala Thr Leu Lys Thr Ile Arg Asn Gly Val His Lys Ile Asp Thr Tyr  
35 40 45ctg aac gcc gcc ttg gac ctc ctg gga ggc gag gac ggt ctc tgc cag 312  
Leu Asn Ala Ala Leu Asp Leu Leu Gly Gly Glu Asp Gly Leu Cys Gln  
50 55 60tat aaa tgc agt gac gga tct aag cct ttc cca cgt tat ggt tat aaa 360  
Tyr Lys Cys Ser Asp Gly Ser Lys Pro Phe Pro Arg Tyr Gly Tyr Lys  
65 70 75 80

ccc tcc cca ccg aat gga tgt ggc tct cca ctg ttt ggt gtt cat ctt	408
Pro Ser Pro Pro Asn Gly Cys Gly Ser Pro Leu Phe Gly Val His Leu	
85 90 95	
aac att ggt atc cct tcc ctg aca aag tgt tgc aac caa cac gac agg	456
Asn Ile Gly Ile Pro Ser Leu Thr Lys Cys Cys Asn Gln His Asp Arg	
100 105 110	
tgc tat gag acc tgt ggc aaa agc aag aat gac tgt gat gaa gaa ttc	504
Cys Tyr Glu Thr Cys Gly Lys Ser Lys Asn Asp Cys Asp Glu Glu Phe	
115 120 125	
cag tat tgc ctc tcc aag atc tgc cga gat gta cag aaa aca cta gga	552
Gln Tyr Cys Leu Ser Lys Ile Cys Arg Asp Val Gln Lys Thr Leu Gly	
130 135 140	
cta act cag cat gtt cag gca tgt gaa aca aca gtg gag ctc ttg ttt	600
Leu Thr Gln His Val Gln Ala Cys Glu Thr Thr Val Glu Leu Leu Phe	
145 150 155 160	
gac agt gtt ata cat tta ggt tgt aaa cca tat ctg gac agc caa cga	648
Asp Ser Val Ile His Leu Gly Cys Lys Pro Tyr Leu Asp Ser Gln Arg	
165 170 175	
gcc gca tgc agg tgt cat tat gaa gaa aaa act gat ctt taa	690
Ala Ala Cys Arg Cys His Tyr Glu Glu Lys Thr Asp Leu	
180 185	
aggagatgcc gacagctagt gacaga	716

<210> 2  
 <211> 189  
 <212> PRT  
 <213> Homo sapiens

<400> 2  
 Met Ala Leu Leu Ser Arg Pro Ala Leu Thr Leu Leu Leu Leu Leu Met  
 1 5 10 15

Ala Ala Val Val Arg Cys Gln Glu Gln Ala Gln Thr Thr Asp Trp Arg  
 20 25 30

Ala Thr Leu Lys Thr Ile Arg Asn Gly Val His Lys Ile Asp Thr Tyr  
 35 40 45

Leu Asn Ala Ala Leu Asp Leu Leu Gly Gly Glu Asp Gly Leu Cys Gln  
 50 55 60

Tyr Lys Cys Ser Asp Gly Ser Lys Pro Phe Pro Arg Tyr Gly Tyr Lys  
 65 70 75 80

Pro Ser Pro Pro Asn Gly Cys Gly Ser Pro Leu Phe Gly Val His Leu  
 85 90 95

Asn Ile Gly Ile Pro Ser Leu Thr Lys Cys Cys Asn Gln His Asp Arg  
 100 105 110

Cys Tyr Glu Thr Cys Gly Lys Ser Lys Asn Asp Cys Asp Glu Glu Phe  
 115 120 125

Gln Tyr Cys Leu Ser Lys Ile Cys Arg Asp Val Gln Lys Thr Leu Gly  
 130 135 140

Leu Thr Gln His Val Gln Ala Cys Glu Thr Thr Val Glu Leu Leu Phe  
 145 150 155 160

Asp Ser Val Ile His Leu Gly Cys Lys Pro Tyr Leu Asp Ser Gln Arg  
 165 170 175

Ala Ala Cys Arg Cys His Tyr Glu Glu Lys Thr Asp Leu  
 180 185

<210> 3

<211> 35

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 3

tttgccggccg catatgggagc tggtctgctgc caagt

35

<210> 4

<211> 37

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 4

tttaagcttc tagaatctgt cactagctgt cggcatc

37

<210> 5

<211> 42

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 5

tttggatcca tcgaaggtcg tcaggagcag gccagaccg ac

42

<210> 6

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 6

gcctttccca cgttatggtt

20

<210> 7

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 7

ggatgtggct ctccactgtt

20

<210> 8

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 8

Gly Cys Gly Ser Pro

1

5

<210> 9

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Consensus  
sequence

<220>

<221> MOD\_RES

<222> (3)..(4)

<223> Any amino acid

<220>

<221> MOD\_RES

<222> (7)

<223> Any amino acid

<400> 9

Cys Cys Xaa Xaa His Asp Xaa Cys

1

5

<210> 10

<211> 182

&lt;212&gt; PRT

&lt;213&gt; Murine sp.

&lt;400&gt; 10

Ser Pro Ala Leu Leu Leu Leu Leu Leu Ala Thr Ala Arg Gly Gln  
 1 5 10 15  
 Glu Gln Asp Gln Thr Thr Asp Trp Arg Ala Thr Leu Lys Thr Ile Arg  
 20 25 30  
 Asn Gly Ile His Lys Ile Asp Thr Tyr Leu Asn Ala Ala Leu Asp Leu  
 35 40 45  
 Leu Gly Gly Glu Asp Gly Leu Cys Gln Tyr Lys Cys Ser Asp Gly Ser  
 50 55 60  
 Lys Pro Val Pro Arg Tyr Gly Tyr Lys Pro Ser Pro Pro Asn Gly Cys  
 65 70 75 80  
 Gly Ser Pro Leu Phe Gly Val His Leu Asn Ile Gly Ile Pro Ser Leu  
 85 90 95  
 Thr Lys Cys Cys Asn Gln His Asp Arg Cys Tyr Glu Thr Cys Gly Lys  
 100 105 110  
 Ser Lys Asn Asp Cys Asp Glu Glu Phe Gln Tyr Cys Leu Ser Lys Ile  
 115 120 125  
 Cys Arg Asp Val Gln Lys Thr Leu Gly Leu Ser Gln Asn Val Gln Ala  
 130 135 140  
 Cys Glu Thr Thr Val Glu Leu Leu Phe Asp Ser Val Ile His Leu Gly  
 145 150 155 160  
 Cys Lys Pro Tyr Leu Asp Ser Gln Arg Ala Ala Cys Trp Cys Arg Tyr  
 165 170 175  
 Glu Glu Ile Thr Asp Leu  
 180

&lt;210&gt; 11

&lt;211&gt; 165

&lt;212&gt; PRT

&lt;213&gt; Rattus sp.

&lt;400&gt; 11

Gln Asp Gln Thr Thr Asp Trp Arg Ala Thr Leu Lys Thr Ile Arg Asn  
 1 5 10 15  
 Gly Ile His Lys Ile Asp Thr Tyr Leu Asn Ala Ala Leu Asp Leu Leu  
 20 25 30  
 Gly Gly Glu Asp Gly Leu Cys Gln Tyr Lys Cys Ser Asp Gly Ser Lys  
 35 40 45  
 Pro Ala Pro Arg Tyr Gly Tyr Lys Pro Ser Pro Pro Asn Gly Cys Gly  
 50 55 60

Ser Pro Leu Phe Gly Val His Leu Asn Ile Gly Ile Pro Ser Leu Thr  
65 70 75 80

Lys Cys Cys Asn Gln His Asp Arg Cys Tyr Glu Thr Cys Gly Lys Gly  
85 90 95

Lys Asn Asp Cys Asp Glu Glu Phe Gln Ser Cys Leu Ser Lys Ile Cys  
100 105 110

Arg Asp Val Gln Lys Thr Leu Gly Leu Ser Gln Asn Val Gln Ala Cys  
115 120 125

Glu Thr Thr Val Glu Leu Leu Phe Asp Ser Val Ile His Leu Gly Cys  
130 135 140

Lys Pro Tyr Leu Asp Ser Gln Arg Ala Ala Cys Trp Cys Arg Tyr Glu  
145 150 155 160

Glu Lys Thr Asp Leu  
165

<210> 12

<211> 136

<212> PRT

<213> Bovine sp.

<400> 12

Asn Ala Ala Leu Asp Leu Leu Gly Gly Glu Asp Gly Leu Cys Gln Tyr  
1 5 10 15

Lys Cys Ser Asp Gly Ser Lys Pro Phe Pro Arg Tyr Gly Tyr Lys Pro  
20 25 30

Ser Pro Pro Asn Gly Cys Gly Ser Pro Leu Phe Gly Val His Leu Asn  
35 40 45

Ile Gly Ile Pro Ser Leu Thr Lys Cys Cys Asn Gln His Asp Arg Cys  
50 55 60

Tyr Glu Thr Cys Gly Lys Ser Lys Asn Asp Cys Asp Glu Ala Phe Gln  
65 70 75 80

Ser Cys Leu Ser Lys Ile Cys Arg Asp Val Gln Lys Thr Leu Gly Leu  
85 90 95

Ala Gln His Val Gln Ala Cys Glu Thr Thr Val Glu Leu Leu Phe Asp  
100 105 110

Ser Val Ile His Leu Gly Cys Lys Pro Tyr Leu Asp Ser Gln Arg Ala  
115 120 125

Ala Cys Arg Cys Arg Tyr Glu Glu  
130 135

<210> 13  
 <211> 194  
 <212> PRT  
 <213> Xenopus sp.

<400> 13  
 Met Arg Phe Arg Gly Phe Leu Tyr Val Leu Trp Phe Ala Tyr Cys Ala  
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 Pro Arg Phe Ser His Gln Glu Pro Trp His Gln Ser Asp Gln Gln Pro  
                   20                  25                  30  
 Glu Thr Pro Asp Trp Arg Met Thr Leu Lys Thr Ile Arg Asn Gly Val  
           35                  40                  45  
 His Lys Ile Asp Met Tyr Leu Asn Ala Ala Leu Asp Leu Leu Gly Gly  
           50                  55                  60  
 Ala Asp Gly Leu Cys His Tyr Glu Cys Arg Asp Gly Ser Lys Pro Val  
           65                  70                  75                  80  
 Pro Arg Tyr Gly Tyr Arg Pro Ser Pro Pro Asn Gly Cys Gly Ser Pro  
                   85                  90                  95  
 Val Phe Gly Val His Asp Ile Gly Ile Pro Ser Met Thr Lys Cys Cys  
           100                  105                  110  
 Asn Gln His Asp Arg Cys Tyr Asp Ser Cys Gly Ile Met Lys Asn Asp  
           115                  120                  125  
 Cys Asp Glu Glu Phe Gln Asn Cys Leu Ser Lys Ile Cys Arg Asp Val  
           130                  135                  140  
 Gln Lys Thr Leu Gly Ile Ser Glu Thr Val Gln Ala Cys Glu Thr Thr  
           145                  150                  155                  160  
 Val Gly Leu Leu Phe Asp Ala Val Ile His Leu Gly Cys Lys Pro Tyr  
                   165                  170                  175  
 Leu Glu Ser Gln Arg Ala Ala Cys Ile Cys Gln Tyr Glu Glu Lys Ile  
           180                  185                  190  
 Asp Leu

<210> 14  
 <211> 37  
 <212> PRT  
 <213> Homo sapiens

<400> 14  
 Glu Tyr Asn Asn Tyr Gly Cys Tyr Cys Gly Leu Gly Gly Ser Gly Thr  
           1                  5                  10                  15  
 Pro Val Asp Glu Leu Asp Lys Cys Cys Gln Thr His Asp Asn Cys Tyr  
                   20                  25                  30

Asp Gln Ala Lys Lys  
35

<210> 15  
<211> 43  
<212> PRT  
<213> Homo sapiens

<400> 15  
Trp Thr Met Pro Gly Thr Leu Trp Cys Gly Val Gly Asp Ser Ala Gly  
1 5 10 15

Asn Ser Ser Glu Leu Gly Val Phe Gln Gly Pro Asp Leu Cys Cys Arg  
20 25 30

Glu His Asp Arg Cys Pro Gln Asn Ile Ser Pro  
35 40

<210> 16  
<211> 38  
<212> PRT  
<213> Conus magus

<220>  
<221> MOD\_RES  
<222> (15)  
<223> Any amino acid

<220>  
<221> MOD\_RES  
<222> (21)  
<223> Any amino acid

<400> 16  
Leu Cys Lys Ile Asn Ser Asn Ala Cys Ser Val Pro Phe Ser Xaa Ile  
1 5 10 15

Pro Cys Gln Lys Xaa Phe Leu Ala Ala Cys Asp Arg His Asp Thr Cys  
20 25 30

Tyr His Cys Gly Lys His  
35

<210> 17  
<211> 41  
<212> PRT  
<213> Oryza sativa

<400> 17  
Pro Leu Leu Arg Tyr Gly Lys Tyr Cys Gly Ile Leu Tyr Ser Gly Cys  
1 5 10 15

Pro Gly Glu Arg Pro Cys Asp Ala Leu Asp Ala Cys Cys Met Val His  
20 25 30



Asp His Cys Val Asp Thr His Asn Asp  
           35                          40

<210> 18

<211> 41

<212> PRT

<213> Homo sapiens

<400> 18

Tyr Lys Pro Ser Pro Asn Gly Cys Gly Ser Pro Leu Phe Gly Val  
   1                  5                  10                  15

His Leu Asn Ile Gly Ile Pro Ser Leu Thr Lys Cys Cys Asn Gln His  
           20                  25                  30

Asp Arg Cys Tyr Glu Thr Cys Gly Lys  
           35                  40